

# Nonlinear Dynamics of a Metabolic Process of the Atherosclerosis

Valeriy Grytsay<sup>1\*</sup>

## Abstract

*A mathematical model of the metabolic process of atherosclerosis is constructed. The functioning of the polyenzymatic prostacyclin-thromboxane system of blood and the influence of a level of "bad cholesterol", namely low density lipoproteins (LDL), on it are studied. With the help of the numerical experiment, we analyze the influence of the concentration of molecules of fat on hemostasis of blood vessels. The kinetic curves for components of the system, phase-periodic bifurcation diagrams, attractors for various modes, and Poincaré cross-section and image of a strange attractor are constructed. The complete spectra of Lyapunov's exponents, divergencies, KS-entropies, predictability horizons, and Lyapunov dimensions of the fractality of strange attractors are calculated. Conclusions about the structural-functional connections, which determine the dependence of hemostasis of a circulatory system on the level of cholesterol in blood are drawn.*

## Keywords

self-organization, hemostasis, chaos, circulatory system, prostacyclin-thromboxane system, Lyapunov's exponents, KS-entropy.

<sup>1</sup> Bogolyubov Institute for Theoretical Physics, National Academy of Sciences of Ukraine, Kyiv, Ukraine

\* [vgrytsay@bitp.kiev.ua](mailto:vgrytsay@bitp.kiev.ua)

## Introduction

In the present work with the help of a mathematical modeling, we continue the study of a prostacyclin-thromboxane system of blood. We will investigate how low density lipoproteins (LDL) influence the dynamics of this metabolic process. In the construction of equations (Eqs.) of our model and the determination of its parameters, we used the results obtained by Prof. S.D. Varfolomeev and Prof. A.T. Mevkh. Their book and the fruitful collaboration with them [1-3] allow Prof. V.P. Gachok and the author to obtain calculation results similar to the experimental ones in the case where the system is in a stable state of hemostasis [4-9], which state characterizes a healthy blood vessel. It is the ideal state, which is attained by synchronization of the systems of thrombosis and antithrombosis. The dynamical stationary equilibrium arises. The desynchronization of these systems results in the appearance of autooscillatory modes in the metabolic process of a prostacyclin-thromboxane system. If the stationary kinetics is broken so that the level of thromboxane increases, then the coagulability of blood grows as well, and the appearance of thrombosis becomes possible in the circulatory system. On the contrary, if the level of prostacyclin increases, then the coagulability of blood decreases, and hemophilia occurs. If the auto oscillatory mode arises, then the appearance of a thrombus as a result of increased coagulability on some time interval and its abruption under a decrease of the coagulability in the following time interval are possible. The actions of external and internal factors induce various modes in the system.

The metabolic process coagulability of blood circulation is considered by the author of the article as an open nonlinear system. The study was conducted using methods of nonlinear dynamics. We will modify the given model by adding four nonlinear Eqs. The other parameters remain unmodified. Within the model, we will study the influence of concentration of "bad cholesterol" (LDL) on the metabolism of a hemostasis of blood vessels. The principal reason for its elevated levels is high dietary fat content. The excessive content of fat in organism causes formation of atheromatous plaques. They are aggregates of LDL on internal walls of blood vessels, causing stenosis. The atheromatous plaques grow over time. As a result, blood circulation slows down, which

creates a deficit of nutrients in tissues. In this case, the arteries become denser and gradually lose their elasticity; i.e., atherosclerosis develops [10].

### 1. The Mathematical Model and Methods of its Study

The general scheme of the hemostasis with regard for the entry of “bad cholesterol” into blood is presented in Fig. 1 [6,10]. According to this scheme, we construct the mathematical model of the given metabolic process (1)-(12) [6,10]:

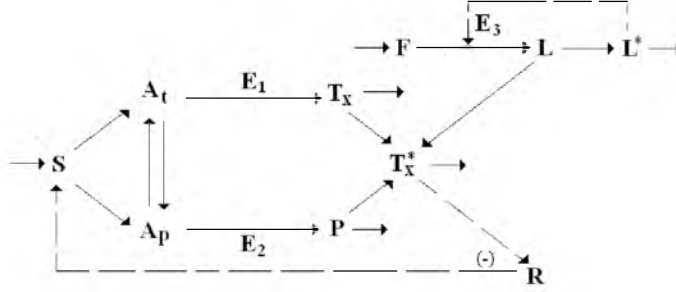


Figure 1. General kinetic scheme of hemostasis

$$\frac{dA_t}{dt} = \frac{k_5 S}{(1+S+R^2)(1+k_6 T_x)} - \frac{k_7 A_t E_1}{(1+A_t+k_1 T_x)(1+E_1)} + k_p A_p - k_t A_t - \alpha_1 A_t, \quad (1)$$

$$\frac{dT_x}{dt} = \frac{k_7 A_t E_1}{(1+A_t+k_1 T_x)(1+E_1)} - \frac{k_8 T_x^4}{(k_9+T_x^4)} - \alpha_2 T_x, \quad (2)$$

$$\frac{dA_p}{dt} = \frac{k_2 S R^2}{(1+S+k_3 A_p)(k_4+R^2)} - \frac{k_{10} A_p E_2}{(1+A_p)(1+E_2)} + k_t A_t - k_p A_p - \alpha_3 A_p, \quad (3)$$

$$\frac{dP}{dt} = \frac{k_{10} A_p E_2}{(1+A_p)(1+E_2)} - \frac{k_{11} T_x^* P^4}{(1+T_x^*)(k_{12}+P^4)} - \alpha_4 P, \quad (4)$$

$$\frac{dE_1}{dt} = \frac{k_{13} A_t}{(1+A_t)(1+R^4)} - \frac{k_7 A_t E_1}{(1+A_t+k_1 T_x)(1+E_1)} - \alpha_5 E_1, \quad (5)$$

$$\frac{dE_2}{dt} = \frac{k_{15} A_p T_x^*}{(k_{16}+A_p)(k_{17}+T_x^*)} - \frac{k_{10} A_p E_2}{(1+A_p)(1+E_2)} - \alpha_6 E_2, \quad (6)$$

$$\frac{dR}{dt} = k_{18} \frac{k_{19}+T_x^*}{k_{20}+(T_x^*+k_{21}R)^4} - \alpha_7 R, \quad (7)$$

$$\frac{dT_x^*}{dt} = k_8 \frac{L+T_x^4}{k_9+L+T_x^4} - \frac{k_{11} T_x^* P^4}{(1+T_x^*)(k_{12}+P^4)} - \alpha_8 T_x^*, \quad (8)$$

$$\frac{dF}{dt} = F_0 - l \frac{E_3}{1+E_3} \cdot \frac{F}{1+F+L}, \quad (9)$$

$$\frac{dL}{dt} = k \frac{E_3}{1+E_3} \cdot \frac{F}{1+F+L} - \mu \frac{LL^*}{1+F+L} - \mu_0 L^*, \quad (10)$$

$$\frac{dL^*}{dt} = \mu_1 \frac{LL^*}{1+L+L^*} - \eta_0 L^* \quad (11)$$

$$\frac{dE_3}{dt} = E_{3_0} L^* \frac{F}{1+F} \cdot \frac{N}{N+L} - \alpha_9 E_3. \quad (12)$$

The model includes the following collection of parameters [6]:

$k = 4$ ;  $k_1 = 3$ ;  $k_2 = 1$ ;  $k_3 = 5$ ;  $k_4 = 10$ ;  $k_5 = 2,1$ ;  $k_6 = 5$ ;  $k_7 = 2$ ;  $k_8 = 1,5$ ;  $k_9 = 5$ ;  $k_{10} = 0,75$ ;

$k_{11} = 0,3$ ;  $k_{12} = 15$ ;  $k_{13} = 0,75$ ;  $k_{15} = 1$ ;  $k_{16} = 0,5$ ;  $k_{17} = 5$ ;  $k_{18} = 5$ ;  $k_{19} = 0,02$ ;  $k_{20} = 25$ ;  $k_{21} = 0,5$ ;  $k_p = 0,1$ ;  $k_t = 0,1$ ;  $S = 2$ ;  $\alpha_1 = 0,01$ ;  $\alpha_2 = 0,01$ ;  $\alpha_3 = 0,01$ ;  $\alpha_4 = 0,173$ ;  $\alpha_5 = 0,05$ ;  $\alpha_6 = 0,07$ ;  $\alpha_7 = 0,2$ ;  $\alpha_8 = 0,0021$ ;  $\alpha_9 = 0,2$ ;  $F_0 = 0,01$ ;  $l = 2$ ;  $\mu = 4$ ;  $\mu_0 = 0,437$ ;  $\mu_1 = 2,3$ ;  $E_{3_0} = 11$ ;  $N = 0,05$ .

The parameters of the system and time are dimensionless quantities [8].

These Eqs. (1)-(12) describe changes of concentrations of the dimensionless corresponding agents.

The input substances in a blood vessel are arachidonic acid  $S$  and molecules of fat  $F$ , which are supplied into blood from the intestinal tract. The output agents of the system are aggregated thrombocytes  $T_x^*$  and oxidized lipoproteins  $L^*$ , which are accumulated on internal walls of arteries. In the model we utilize the law of mass action and the kinetics of enzyme catalysis. The Eqs. involve the balance of masses of the intermediate products of reactions on separate stages of the metabolic process.

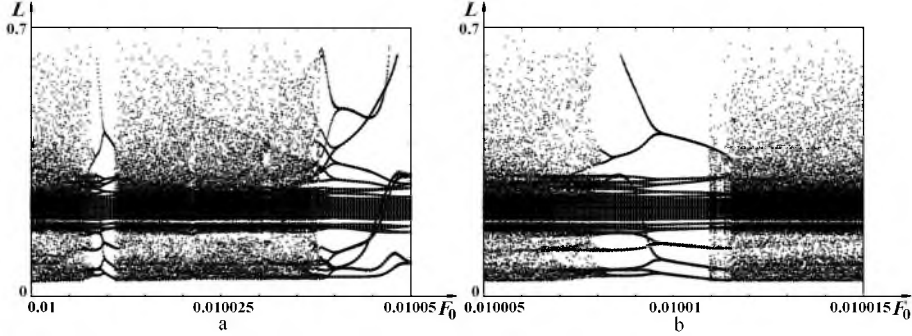
Eqs. (1) and (3) describe, respectively, changes in the concentrations of arachidonic acid  $A_t$  and  $A_p$  in thrombocytes and in endothelial cells of the vessel. These processes are affected by activity of the corresponding phospholipases. The coefficients  $k_5$  and  $k_2$  characterize, respectively, the rate of these processes. The accumulated arachidonic acid is transformed then by prostaglandin- $H$ -synthase of thrombocytes  $E_1$  and prostaglandin- $H$ -synthase of prostacyclins  $E_2$ . Respectively, thromboxanes  $T_x$  (2) and prostacyclins  $P$  (4) are formed. The rate of these enzymatic processes is determined by the coefficients  $k_7$  and  $k_{10}$ . The coefficients  $k_t$  and  $k_p$  characterize the exchange by arachidonic acid between thromboxanes and endothelial cells. Eqs. (5) and (6) describe, respectively, changes of the concentrations of enzymes  $E_1$  and  $E_2$ . The coefficients  $k_{13}$  and  $k_{15}$  determine the intensity of biosynthesis in thromboxanes and endothelial cells. The inactivation of these enzymes is guided by the corresponding terms with the coefficients  $k_7$  and  $k_{10}$  in Eqs. (2) and (4). Eqs. (1)-(6) satisfy completely the balance of masses in enzyme catalysis. Eq. (7) describes changes in the concentration of the controlling component, cyclic adenosine monophosphate ( $cAMP$ )  $R$ . Its presence in Eqs. (1) and (3) creates the negative feedback that affects the level of activity of phospholipases of thrombocytes  $A_t$  and endothelial cells  $A_p$ . Equation (8) describes aggregation of thrombocytes  $T_x^*$  and their dissipation under effect of prostacyclin. Its concentration depends also on blood LDL level. Eqs. (9)-(12) characterize the metabolic process of formation of LDL in blood and the accumulation of plaques on walls of the vessel. Fat molecules  $F$  (9) are supplied by blood to arteries from liver and small intestine. Eqs. (9)-(10) describe the process of creation of "bad cholesterol"  $L$  from fat. Its deposition on walls of blood vessels in the form of oxidized lipoproteids  $L^*$  (plaques) is described by Eqs. (10)-(11). In the metabolic process, the positive feedback controlled by enzyme  $E_3$  is formed (12). The accumulation of cholesterol in arteries and the growth of plaques cause thrombophilia. In this case, the lumen of an artery becomes narrower, i.e., stenosis develops. The above Eqs. involve also the dissipation of the corresponding substances at the expense of other metabolic processes and the flow of blood in an artery.

The given system is an open non equilibrium one. Its study was carried out with the use of the theory of nonlinear differential equations [11-12] and the methods of mathematical modeling used earlier in [13-24].

## 2. Result of Studies

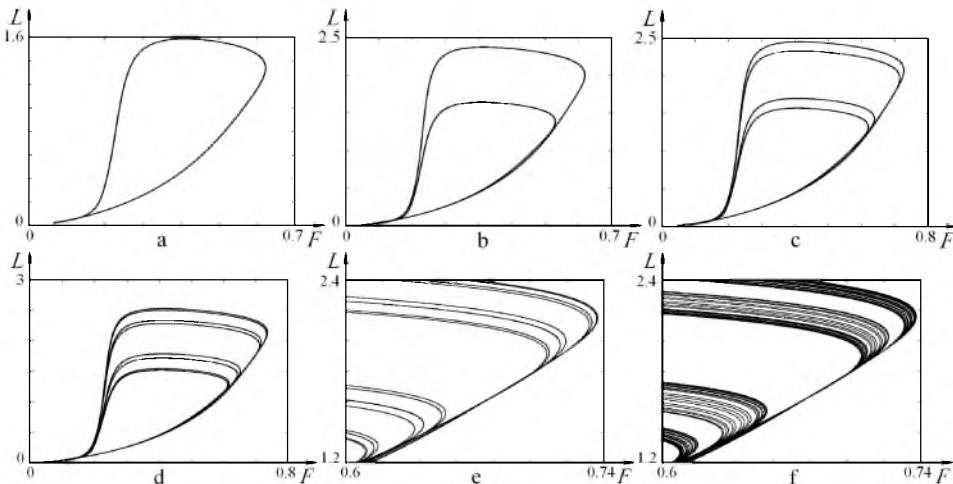
The investigation of the mathematical model (1)-(12) has shown that, in addition to the ideal stationary modes of the metabolic process of the thrombosis-antithrombosis system [4-9], the model includes also autooscillatory modes. Depending on the rate of supply of molecules of fat  $F_0$  to blood, the level of "bad cholesterol"  $L$  varies. The metabolic process of hemostasis becomes unstable. The study of autooscillatory modes will enable us to comprehend the dynamics of the metabolic process and to reveal the structural-functional connections in this system.

In Fig. 2, we show a phase-parametric diagram of the system for  $L(t)$  at a variation of  $F_0$  in the corresponding intervals. In order to construct the phase-parametric diagrams, we have used the cutting method. In the phase space of a trajectory of the system, we place a cutting plane for the value of concentration of cyclic adenosine monophosphate (c AMP)  $R = 1.77$ . If the trajectory crosses this plane in some direction, we mark the value of chosen variable on the phase-parametric diagram ( $L(t)$  in this case). Such choice is explained by the symmetry of oscillations of the given component relative to such point in multiply calculated earlier modes. For every value of  $L$ , we mark the intersection of the trajectory and this plane after the trajectory falls into the attractor. If a multifold periodic limiting cycle arises, we will see a number of points on the plane, which coincide in the period. If a deterministic chaos arises, the points of intersection are located chaotically.



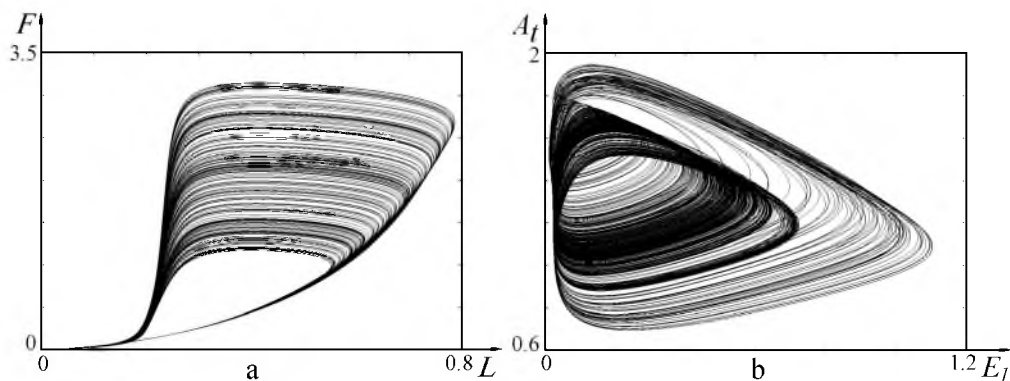
**Figure 2.** Phase-parametric diagram of the system for the variable  $L(t)$  : a -  $F_0 \in (0.01, 0.01005)$  ; b -  $F_0 \in (0.010005, 0.010015)$

It is convenient to consider the phase-parametric diagram from right to left. As  $F_0$  decreases (Fig. 2,a), we observe the successive transition to autoperiodic modes with higher multiplicity due to a cascade of bifurcations with the doubling of a period, until the chaotic mode is finally established due to the intermittence. Such scenario is shown in Fig.3,a-f. The phase portraits of regular attractors transfer to a strange attractor. As the given parameter decreases further, the chaotic modes hold in the system. In the interval  $F_0 \in (0.010007, 0.010012)$  (Fig. 2,a,b), chaos is destroyed, and the periodicity window arises. In this window, the transition from autoperiodic modes to chaotic ones occurs also as a result of the cascade of bifurcations with the doubling of a period by Feigenbaum's scenario. The transition finishes analogously to the previous scenario. Due to the intermittence, chaos is formed. The analogy of these scenarios indicates the fractal nature of the given cascades of bifurcations.



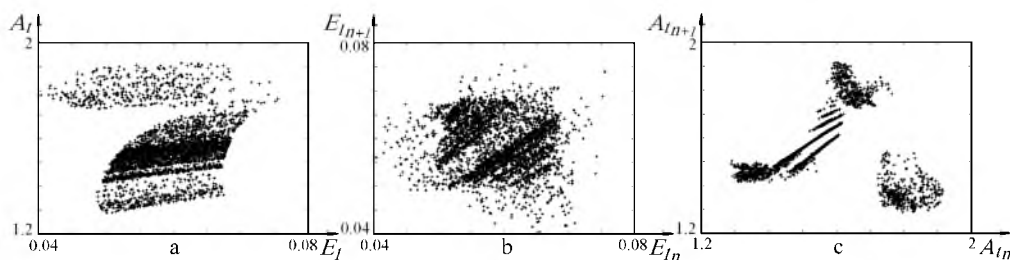
**Figure 3.** Projections of the phase portraits of attractors of the system in the plane  $(F, L)$  : a -  $1 \cdot 2^0$ , for  $F_0 = 0.0102$  ; b -  $1 \cdot 2^1$ , for  $F_0 = 0.01005$  ; c -  $1 \cdot 2^2$ , for  $F_0 = 0.010045$  ; d -  $1 \cdot 2^4$ , for  $F_0 = 0.0100383$  ; e -  $1 \cdot 2^8$ , for  $F_0 = 0.010037$  ; f -  $1 \cdot 2^x$ , for  $F_0 = 0.010036$

In Fig. 4,a,b, as an example, we show the projections of the strange attractor  $2^x$  for  $F_0 = 0.01$  in the planes  $(L, F)$  and  $(E_1, A_1)$ . The obtained strange attractor is formed due to the funnel effect. An element of the phase volume of such attractor is stretched in some direction and contracts in other directions, by preserving its stability. Therefore, the mixing of trajectories happens in narrow contracted regions of the phase space of a funnel, and the deterministic chaos arises.



**Figure 4.** Projections of phase portraits of strange attractor  $2^x$ , for  $F_0 = 0.01$ : a – in the plane  $(L, F)$ , b – in the plane  $(E_1, A_1)$

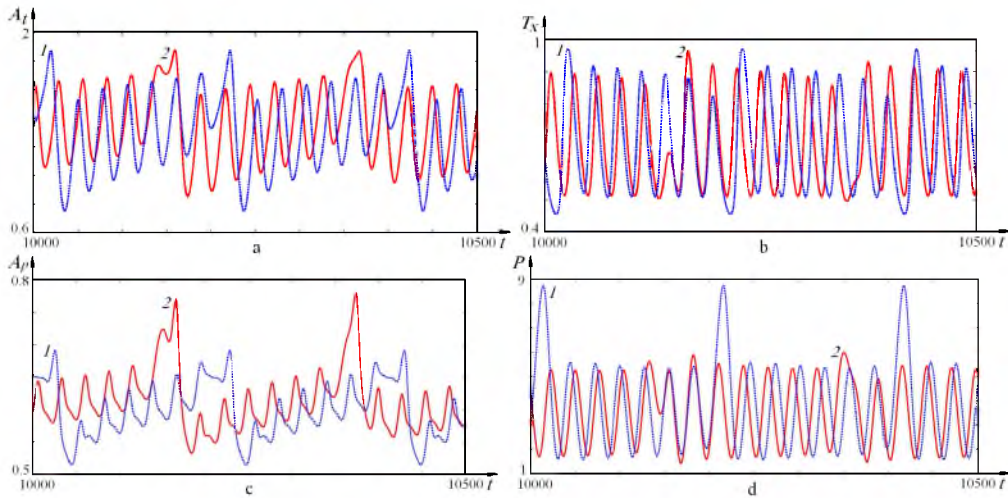
For the given strange attractor in Fig. 5 a,b,c, we constructed the projection of the intersection with the plane  $R = 1.77$  and the Poincaré image. The intersection plane was chosen so that the phase trajectory  $R(t)$  crosses it the maximal number of times, as the given component decreases, without any touching of the intersection plane by the phase curve.



**Figure 5.** Intersection projection (a) and the Poincaré images (b), (c) with the plane  $R = 1.61$  for a strange attractor formed for  $F_0 = 0.01$

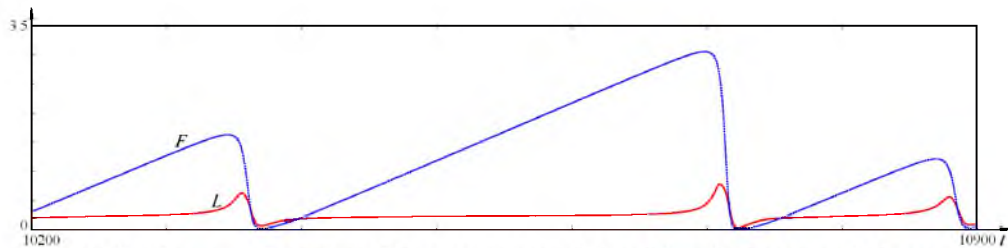
The obtained intersection points and the Poincaré image do not possess a geometric self-similarity. The number of points permanently increases with the duration of a numerical integration of the system. This demonstrates the chaoticity of the attractor and the impossibility of some reduction of the given complicated kinetic scheme of metabolic processes to a one-dimensional discrete approximation of the system under study.

In Fig. 6, we compare the kinetics of some components of the system in the periodic (1) and chaotic (2) modes.



**Figure 6.** Kinetic curves of the metabolic process of atherosclerosis during the running of the autoperiodic (1) ( $F_0 = 0.0102$ ) and chaotic (2) ( $F_0 = 0.01$ ) modes; a -  $A_t$ ; b -  $T_x$ ; c -  $A_p$ ; d -  $P$ .

Changes of the concentrations of fat and “bad cholesterol” in the chaotic metabolic mode of atherosclerosis are shown in Fig. 7. Such nonuniform change of LDL affects the thrombosis-antithrombosis system, by destroying a steady hemostasis of an artery. The balance between the amounts of cholesterol deposited in a blood vessel and that taken out with blood is violated. At certain point the amount of deposited cholesterol becomes larger. This favors the formation of plaques in a blood vessel. Thus, the appearance of atherosclerosis depends on the self-organization of the metabolic process in the thrombosis-antithrombosis system. Under the self-organization, blood vessels adapt to the conditions of nutrition. If a desynchronization of these processes occurs, the risk of atherosclerosis development becomes more significant.



**Figure 7.** Kinetic curves of variations of the concentrations of  $F$  and  $L$  in the chaotic metabolic process of atherosclerosis for  $F_0 = 0.01$

For the unique identification of the type of the obtained attractors and for the determination of their stability for various values of parameter  $F_0$ , we calculated the complete spectra of Lyapunov's exponents and their sum. The calculation was carried out by Benettin's algorithm with the orthogonalization of the vectors of perturbations by the Gram-Schmidt method.

Using the Pesin theorem and the values of Lyapunov's exponents, we calculated also the KS-entropy (Kolmogorov-Sinai entropy)  $h$  and the Lyapunov index “predictability horizon”  $t_{\min}$ . The Lyapunov dimension  $D_{Fr}$  of the fractality of strange attractors was found by the Kaplan-Yorke formula.

Below, as an example for comparison, we present some results of calculations of the mentioned indices.

**Table 1.** Calculations of the mentioned indices

$F_0$	Attractor	$\lambda_1$	$\lambda_2$	$\lambda_3$	$h$	$t_{\min}$	$D_{Fr}$
0.01	$1 \cdot 2^x$	.00192	-.00020	-.00432	.00192	520.83	2.44
0.010005	$1 \cdot 2^x$	.00172	-.00018	-.00370	.00172	581.40	2.46
0.01001	$1 \cdot 2^0$	.00004	.00044	-.00155	-	-	-
0.010015	$1 \cdot 2^x$	.00175	.00013	-.00322	.00175	571.43	2.54
0.01002	$1 \cdot 2^x$	.00150	.00016	-.00296	.00150	666.67	2.51
0.010025	$\approx n \cdot 2^0$	.00033	.00009	-.00173	-	-	-
0.01003	$\approx n \cdot 2^0$	.00097	.00018	-.00388	-	-	-
0.010035	$\approx n \cdot 2^0$	.00065	.00013	-.00408	-	-	-
0.01005	$\approx n \cdot 2^0$	.00002	-.00005	-.00158	-	-	-

These results show the variety of geometric structures of the obtained attractors and the predictability of the metabolic process depending on the concentration of molecules of fat in blood and on the level of “bad cholesterol”.

Calculating successively various strange attractors, we can find some regularity in the hierarchy of their chaotic behavior. Respectively, the variation of the given indices changes a geometric view of attractors of the system.

Autooscillations in the metabolic process of hemostasis of a blood vessel arise due to the interaction between thrombosis and antithrombosis systems of blood, which is regulated by the level of cyclic adenosine monophosphate. The presence of “bad cholesterol” in blood causes desynchronization of these systems and the appearance of chaotic modes in the metabolism of a hemostasis. *LDL* affects the binding of thrombocytes and deposits on the walls of blood vessels. This leads to the autocatalysis of cholesterol in blood.

Thus, the hemostasis under a change of the amount of cholesterol in blood characterizes the adaptation of the metabolic process of a blood vessel to these changes, by preserving its functionality in this case.

## Conclusions

We have constructed a mathematical model of the process of atherosclerosis of a blood vessel. The mathematical model describes the metabolic process of the thrombosis-antithrombosis system based on the prostacyclin-thromboxane system of blood. We have studied how molecules of *LDL* affect the imbalance of this system. The autooscillatory modes determined with this model indicate a complicated internal dynamics of formation of the self-organization in a blood vessel, i.e. that of the hemostasis. We have studied the dependence of autooscillatory modes on the concentration of fat in blood. Moreover, we determined the chaotic modes of strange attractors. During such modes, the imbalance between the amount of “bad cholesterol” deposited in a blood vessel and its removal from the system happens. This provokes the formation of plaques in an artery. It is shown that *LDL* affects the binding of thrombocytes and deposits on walls of blood vessels. This causes the autocatalysis of cholesterol in blood and the increase of its level. The mathematical study of the obtained modes is performed. The phase-parametric diagram, kinetic curves, projection of phase portraits, and Poincaré cross-sections and images are constructed. The Lyapunov's exponents, divergencies, “predictability horizons,” and Lyapunov dimensions of the fractality of strange attractors are calculated. These indices characterize the stability and structure of calculated attractors.

The obtained results clarify the metabolic process of hemostasis and find the structural-functional connections affecting the appearance of atherosclerosis of blood vessels.

## References

- [1] S.D. Varfolomeev, A.T. Mevkh. // *Prostaglandins -- Molecular Bioregulators*. – Moscow University, Moscow, 1985 (in Russian).
- [2] S.D. Varfolomeev, A.T. Mevkh, V.P. Gachok Kinetic model of the multienzyme system of blood prostanoïd synthesis. 1. Mechanism of stabilization of the levels of thromboxane and prostacyclin// *Molek. Bill.* - 1986. - Vol. 20, No.4, - pp.957-966.
- [3] S.D. Varfolomeev, V.P. Gachok, A.T. Mevkh, Kinetic behavior of the multienzyme system of blood prostanoïd synthesis // *BioSystems*. – 1986. – Vol. 19, - P. 45-54.
- [4] V.I. Grytsay, The conditions of the self-organization in the multienzyme prostacyclin-thromboxane system. // *Visn. Kyiv. Univ.*, 2002, No. 3, pp. 372-376.
- [5] V.I. Grytsay, V.P. Gachok, The modes of self-organization in prostacyclin-thromboxane system // *Visn. Kyiv. Univ.*, 2002, No. 4, pp. 365-370.
- [6] V.I. Grytsay, V.P. Gachok, Ordered structures in the mathematical system of prostacyclin and thromboxane model // *Visn. Kyiv. Univ., Ser. Fiz.-Mat. Nauk.*, 2003, No. 1, pp. 338 – 343.
- [7] V.I. Grytsay, Modeling of processes in the multienzyme prostacyclin and thromboxane system // *Visn. Kyiv. Univ.*, 2003, No. 4, pp. 379 – 384.
- [8] V.P. Gachok, Kinetics of Biochemical Processes. – *Naukova Dumka*, Kiev, 1988 (in Russian)
- [9] V.P. Gachok, Strange Attractors in Biosystems– *Naukova Dumka*, Kiev, 1989 (in Russian)
- [10] P. Libby, Atherosclerosis: The New View, Scientific American, on May 1, pp. 46-55 (2002).
- [11] V.S. Anishchenko, *Complex Oscillations in Simple Systems*. Moscow, Nauka, 1990 (in Russian).
- [12] S.P. Kuznetsov, *Dynamical Chaos*. Moscow, Nauka, 2001 (in Russian).
- [13] V.I. Grytsay. Self-organization of biochemical process of immobilized cells bioselective of membrane biosensor, *Ukr. J. Phys.*, Vol. 46, 124-127, 2001.
- [14] V.V. Andreev and V.I. Grytsay. Modeling of nonactive zones in porous granules of a catalyst and in a biosensor. *Matem. Modelir.* vol. 17, no. 2, 57-64, 2005.
- [15] V.V. Andreev and V.I. Grytsay. Influence of heterogeneity of diffusion-reaction process for the formation of structures in the porous medium. *Matem. Modelir.* vol. 17, no. 6, 3 – 12, 2005.
- [16] V.I. Grytsay and V.V. Andreev. The role of diffusion in the active structures formation in porous reaction-diffusion media. *Matem. Modelir.* vol. 18, no. 12, 88-94, 2006.
- [17] V.I. Grytsay. Structural instability of a biochemical process. *Ukr. J. Phys.*, vol. 55, No. 5, 599-606, 2010.
- [18] V.I. Grytsay and I. V. Musatenko. The structure of a chaos of strange attractors within a mathematical model of the metabolism of a cell. *Ukr. J. Phys.* vol. 58, No. 7, 677-686, 2013.
- [19] V.I. Grytsay, I. V. Musatenko, A mathematical model of the metabolism of a cell, Self-organization and Chaos, *Chaotic Modeling and Simulation (CMSIM)*, No. 4, 539-552, 2013.
- [20] V. Grytsay, I. Musatenko, Nonlinear self-organization dynamics of a metabolic process of the Krebs cycle, *Chaotic Modeling and Simulation (CMSIM)*, 2014, Vol. 3, p. 207-220.
- [21] V. Grytsay, Lyapunov indices and the Poincaré mapping in a study of the stability of the Krebs cycle. *Ukr. J. Phys.* Vol 60, No. 6, 564-577, 2015.
- [22] V.I. Grytsay, Self-organization and fractality in the metabolic process of glycolysis, *Ukr. J. Phys.*, 2015, vol. 60, No. 12, 1253-1265.
- [23] V. Grytsay, Self-organization and fractality created by gluconeogenesis in the metabolic process, *Chaotic Modeling and Simulation (CMSIM)*, 2016, Vol. 2, p. 113-127.
- [24] V.I. Grytsay, Self-organization and chaos in the metabolism of hemostasis in a blood vessel, *Ukr. J. Phys.*, 2016, vol. 61, No. 7, 648-655.